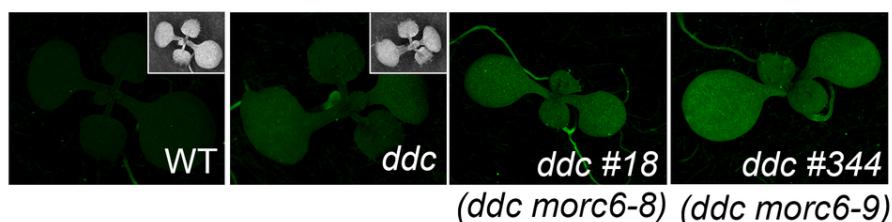
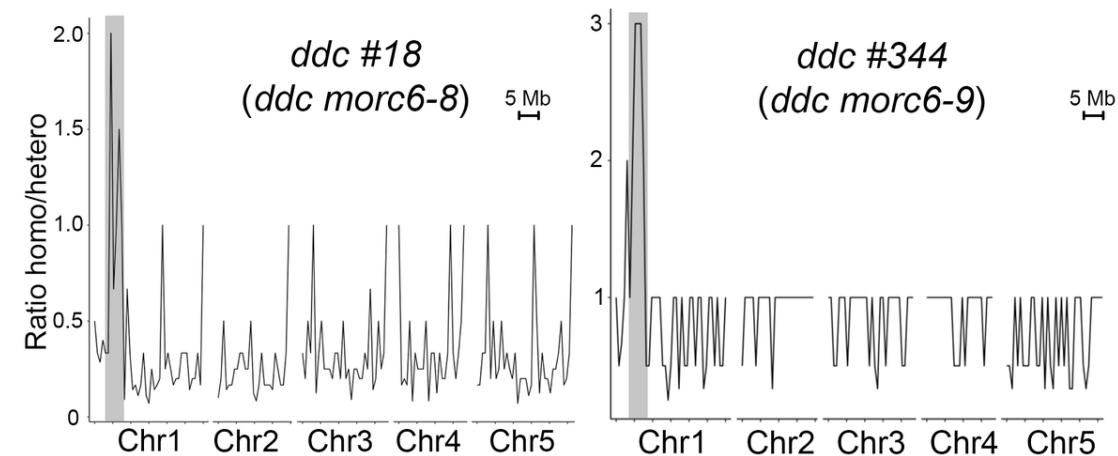


S1 fig

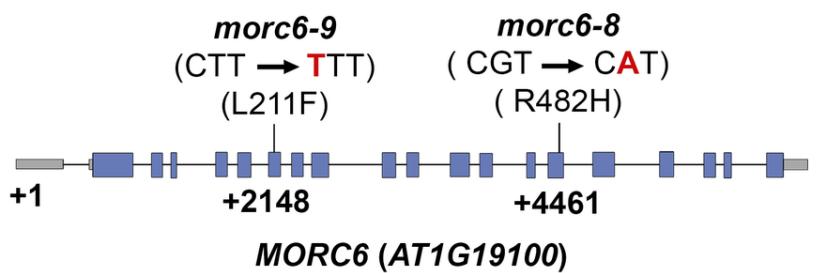
A



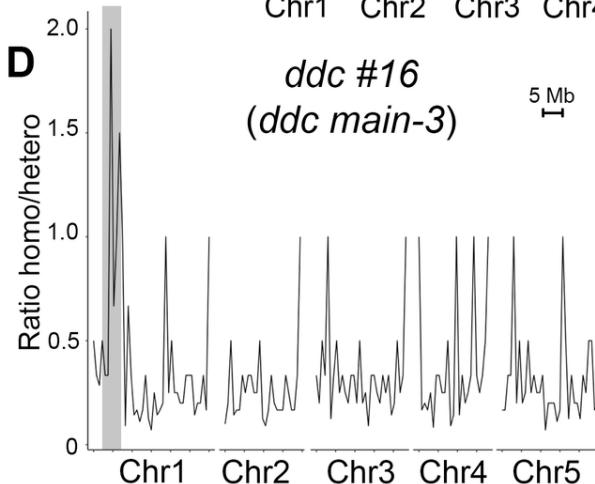
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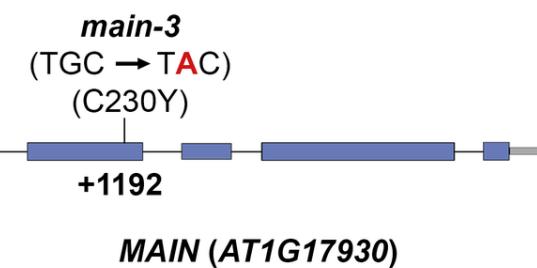
C



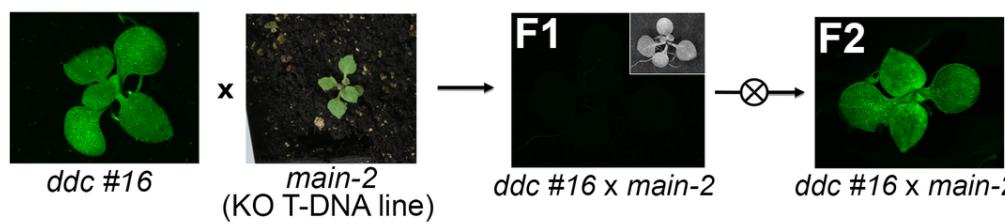
D



E

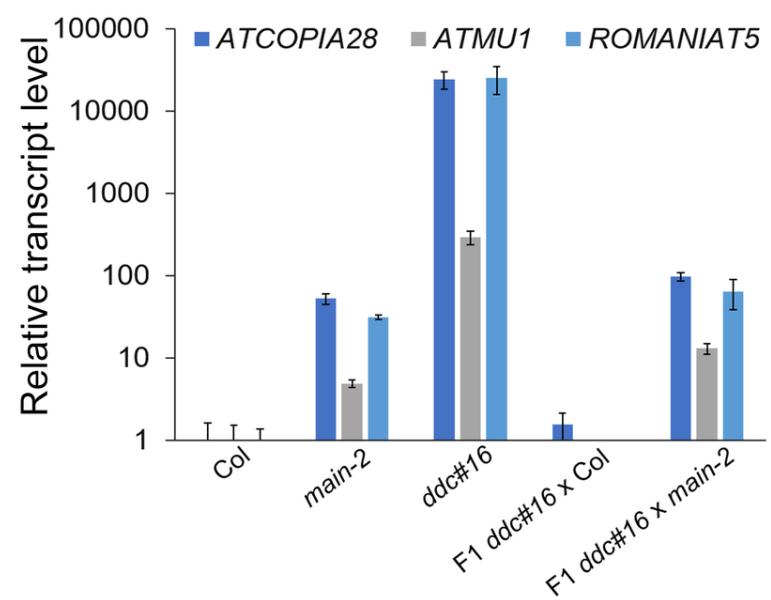


F

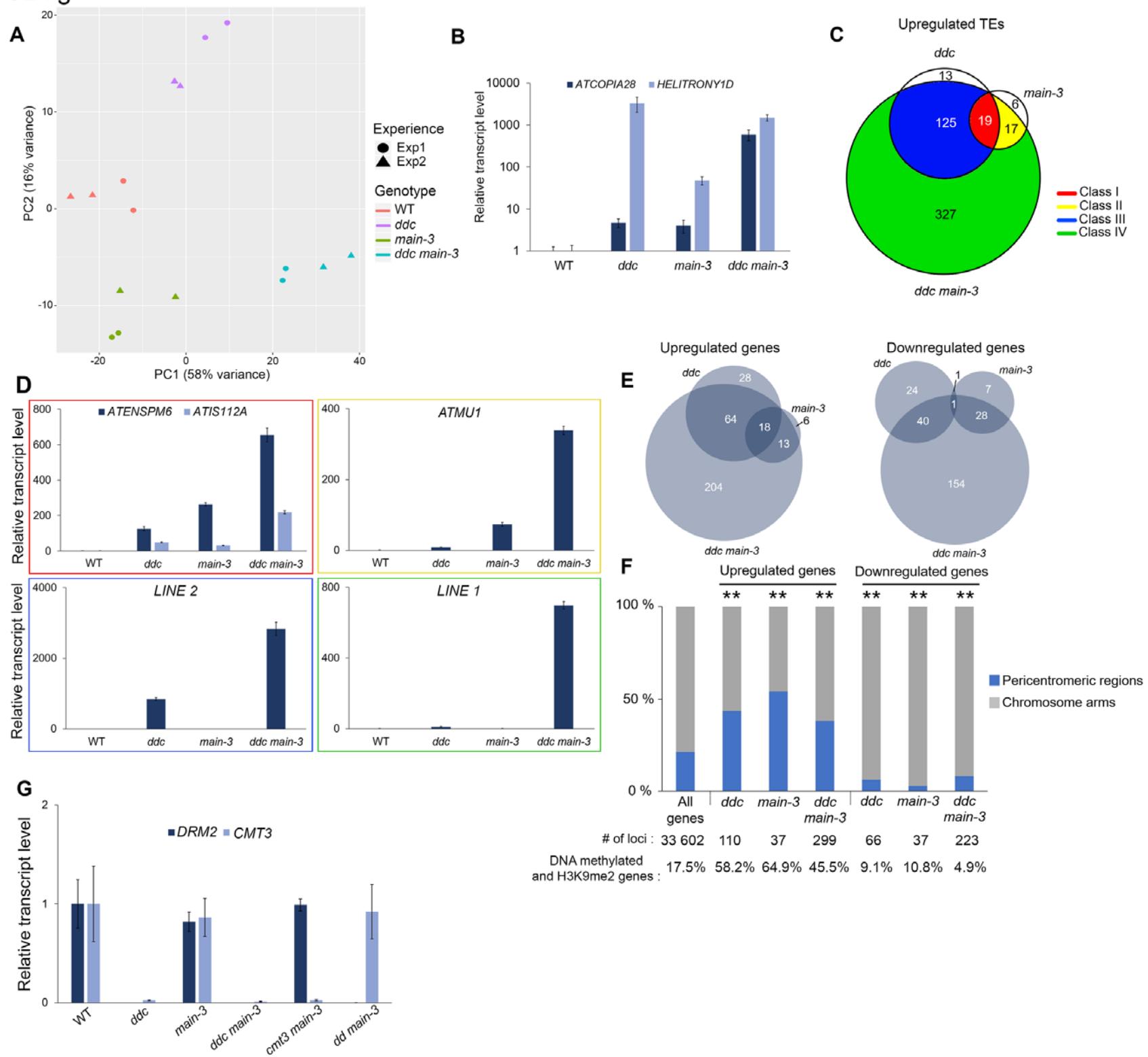


main	F2 plants (n°)									
	1	2	3	4	5	6	7	8	9	10
EMS mutation	Ho	He	Ho	Ho	He	He	He	He	He	He
T-DNA mutation	WT	He	WT	WT	He	He	He	He	He	He
<i>drm2</i>	Ho	Ho	Ho	Ho	Ho	Ho	Ho	Ho	Ho	Ho
<i>cmt3</i>	Ho	He	Ho	Ho	Ho	He	Ho	He	He	He

G

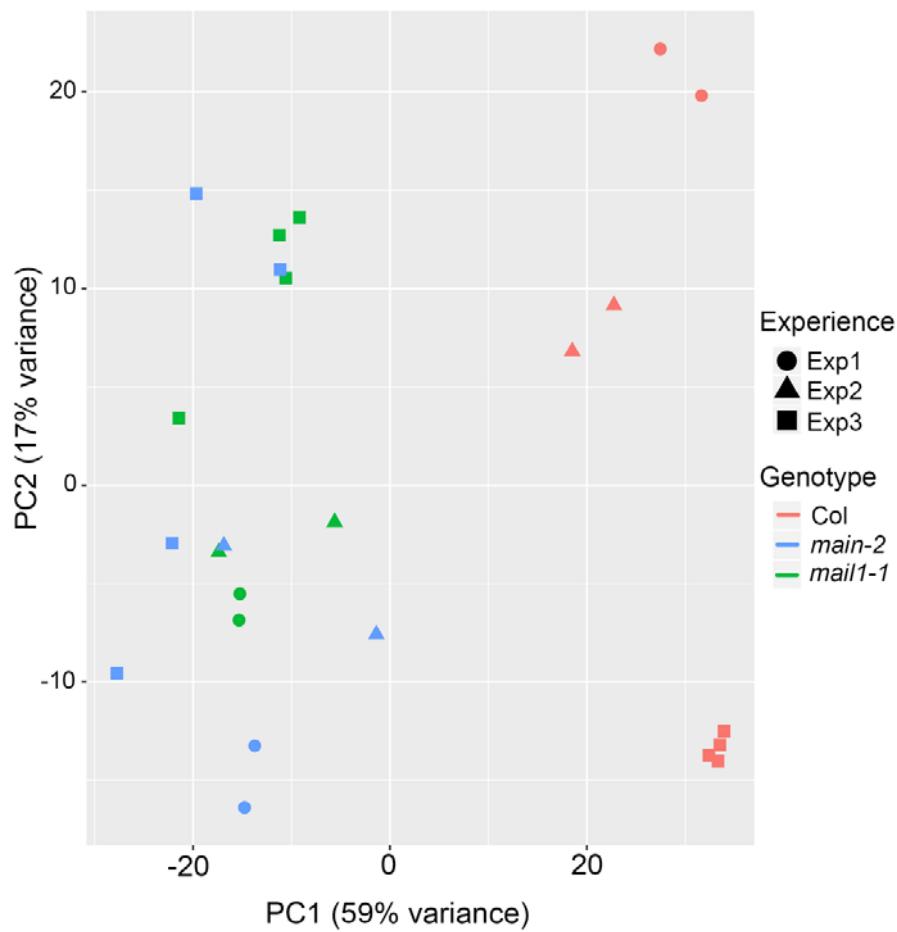


S2 fig

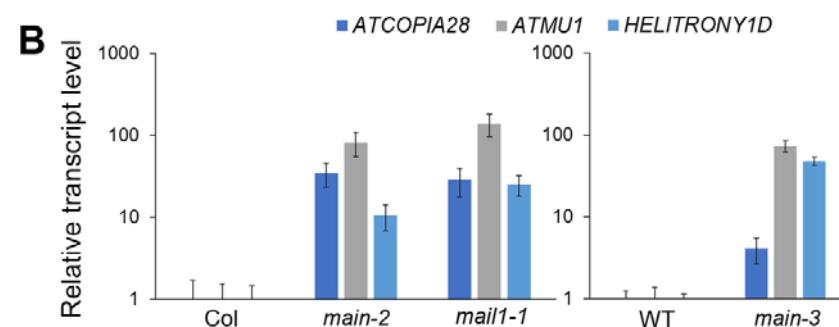


S3 fig

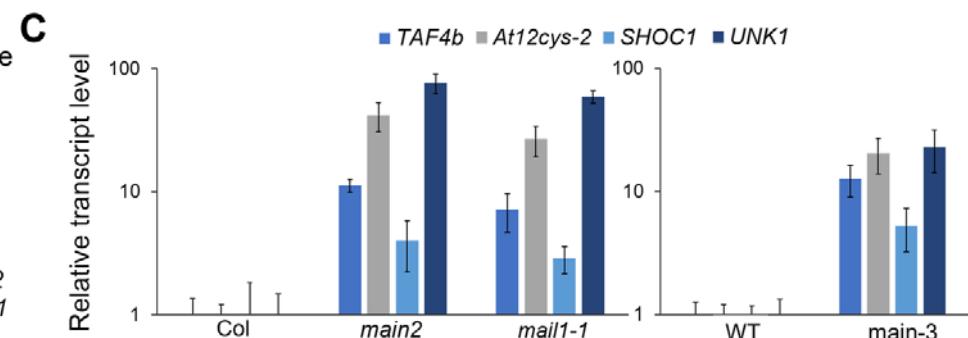
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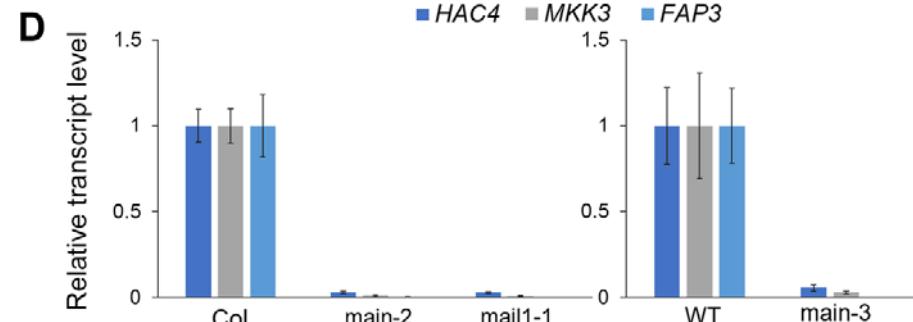
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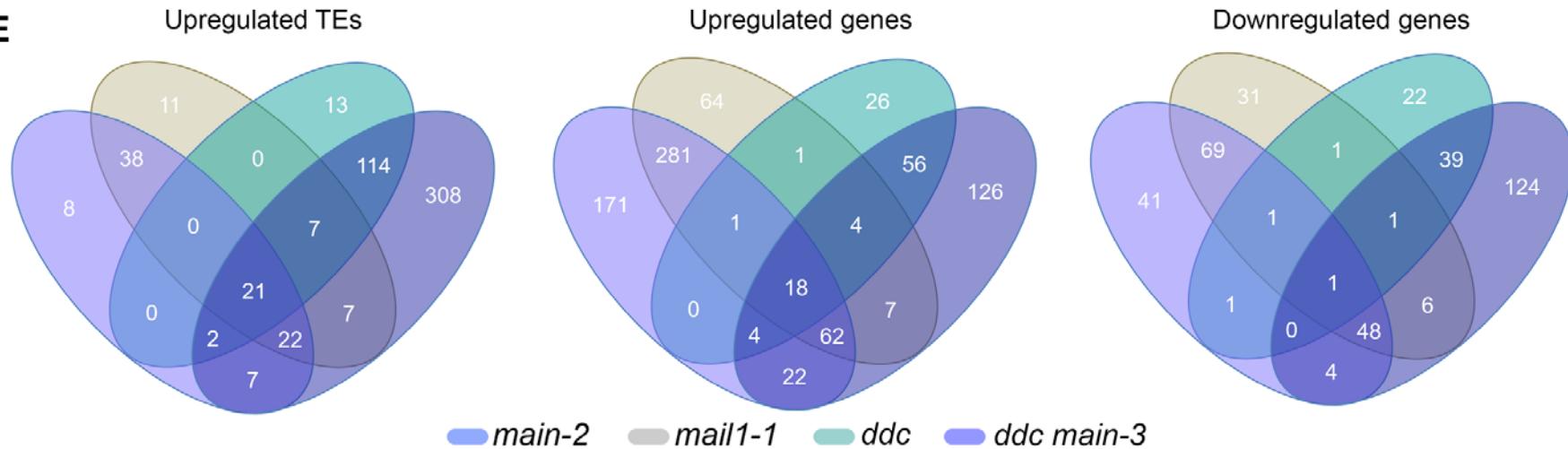
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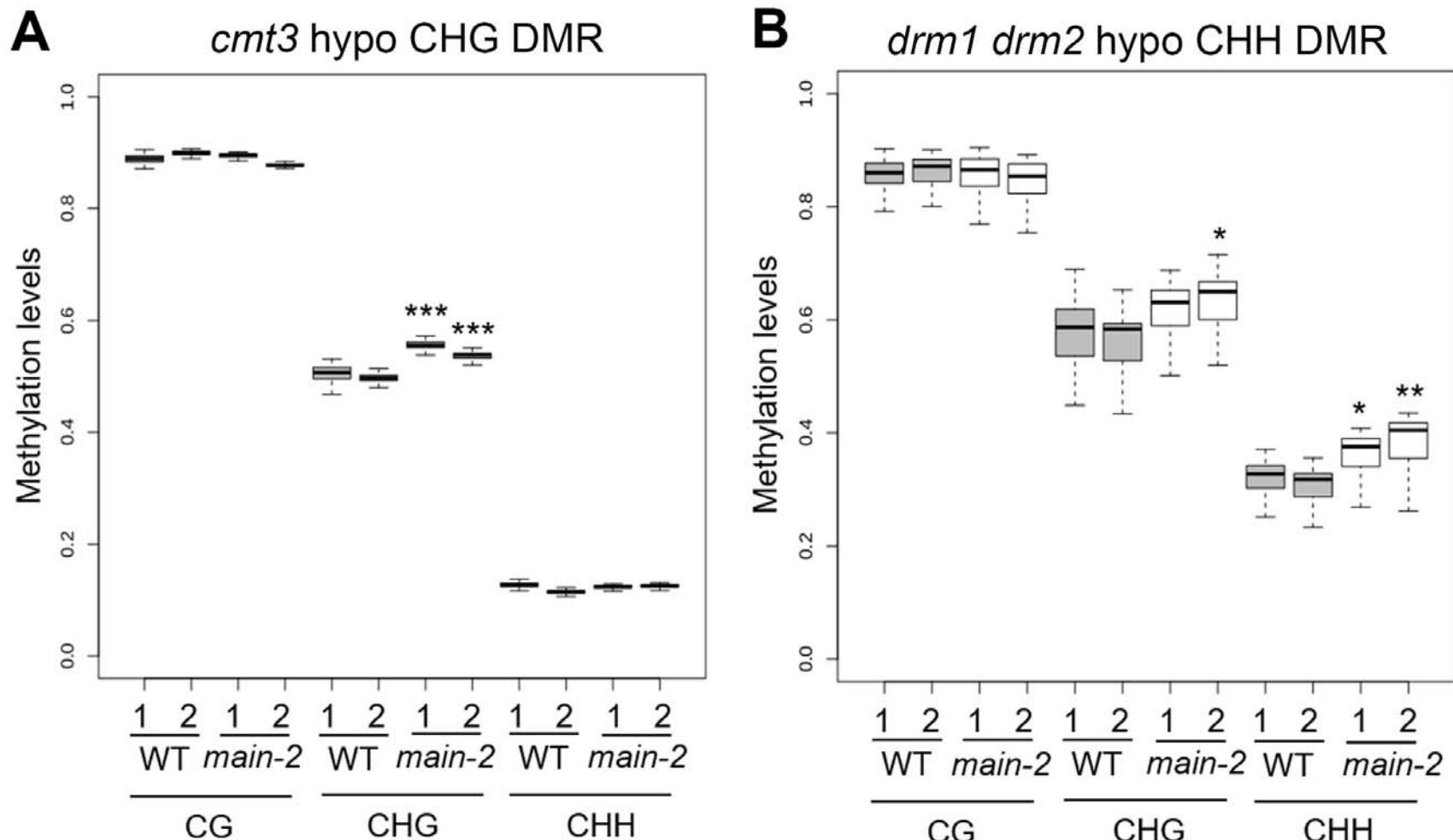
D



E



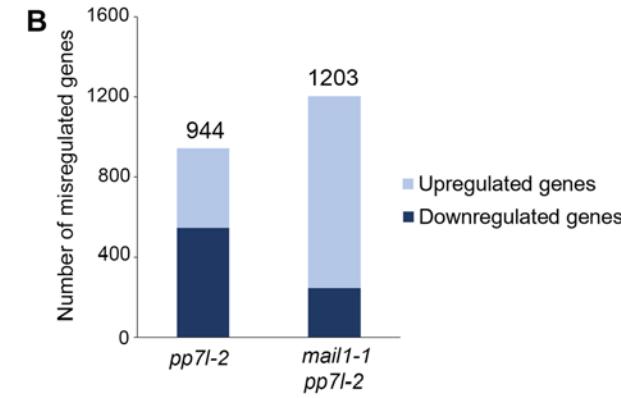
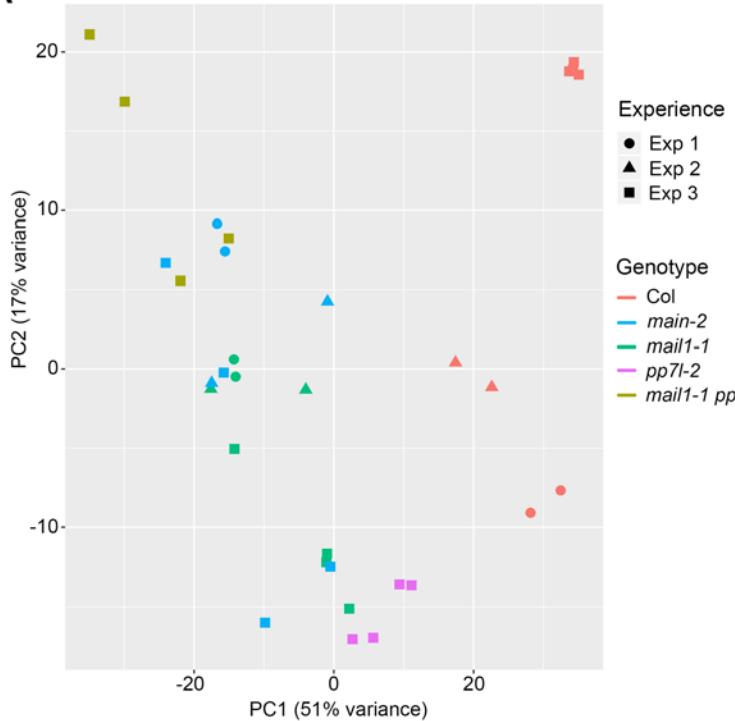
S4 fig



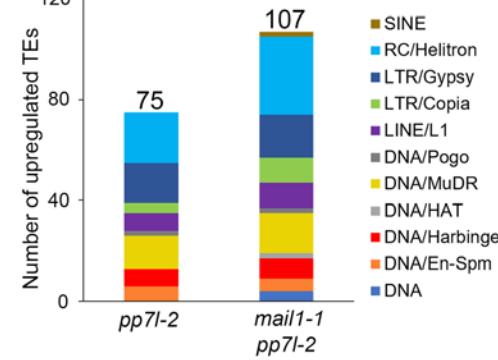
C

			TEs	Genes	Intergenic regions
			CG	CHG	CHH
<i>main-2</i>	Hyper DRM	CG	4	7	0
		CHG	0	0	1
		CHH	0	0	0
	Hypo DMR	CG	48	28	14
		CHG	0	0	0
		CHH	0	0	0

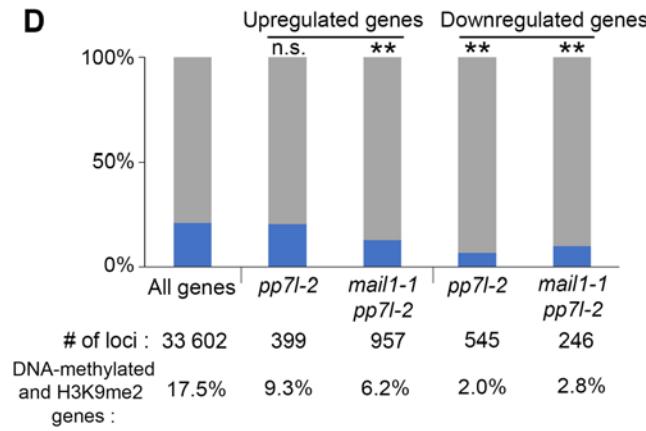
S5 fig
A



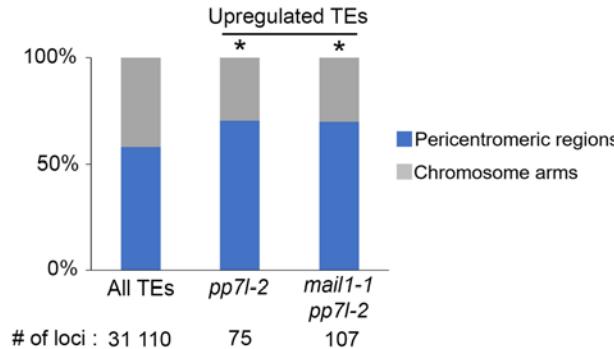
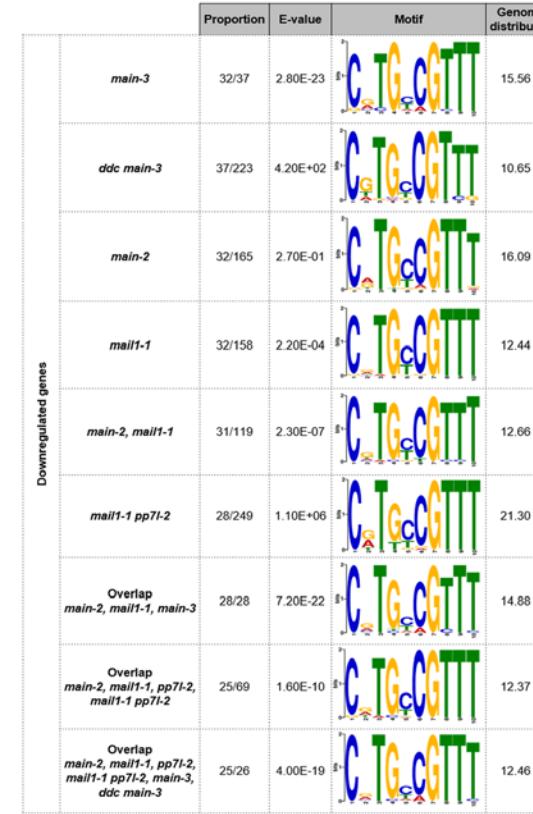
C



D



E



S6 fig

Fig 1C

kDa

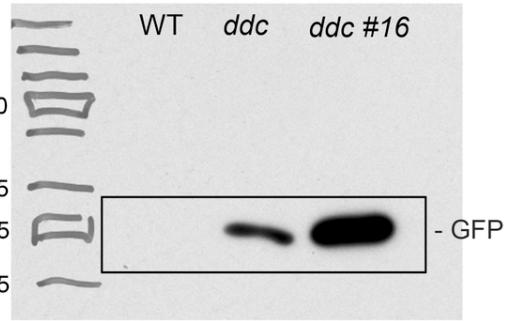


Fig 5B

kDa

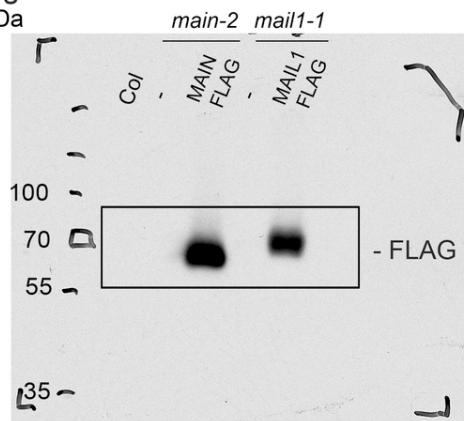


Fig 5B

kDa

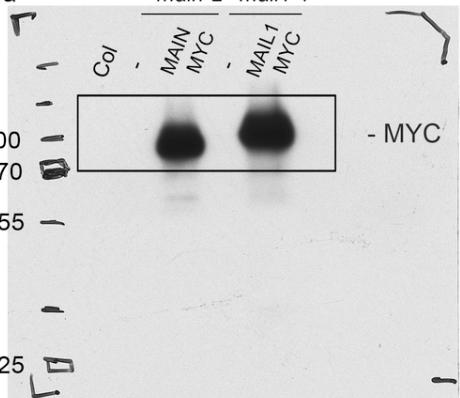


Fig 5G

kDa

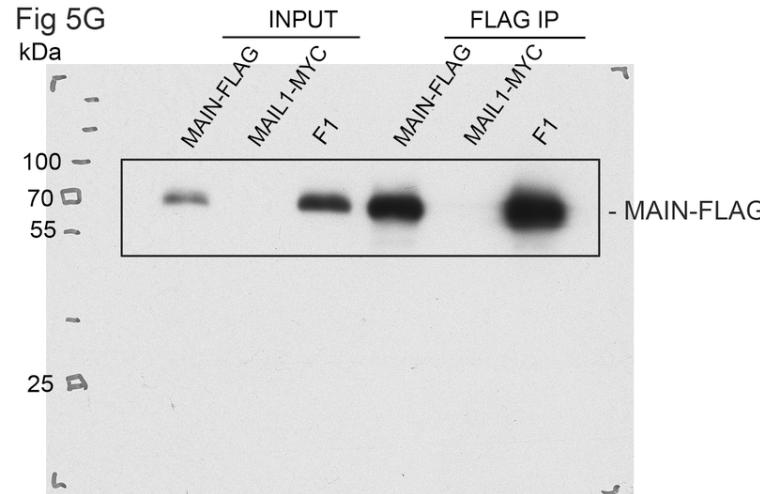


Fig 5G

kDa

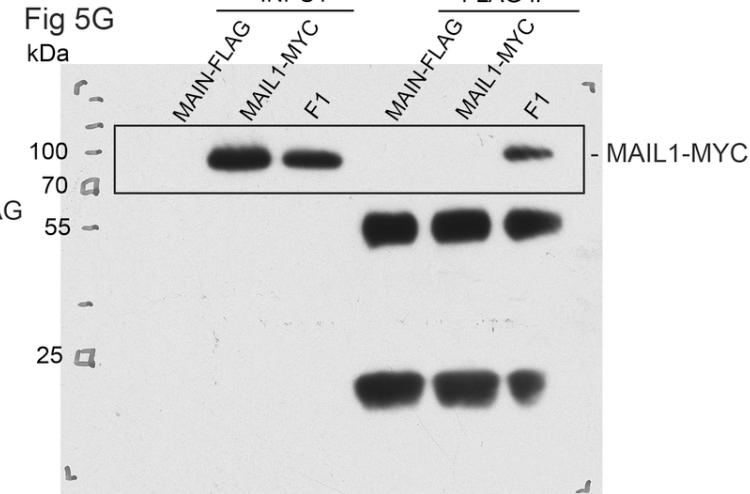


Fig 5H

kDa

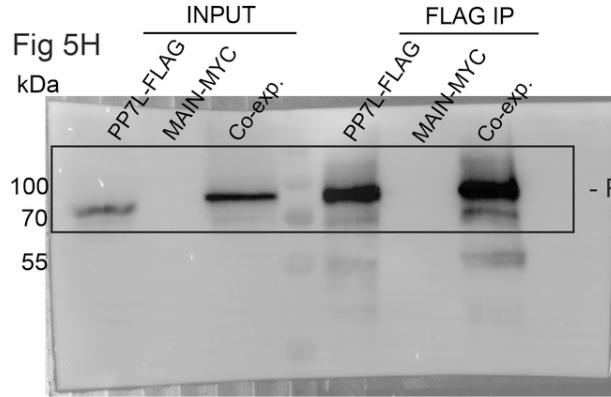


Fig 5H

kDa

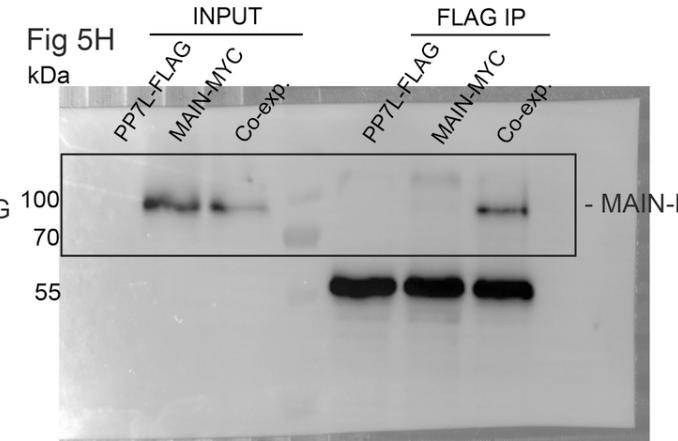


Fig 5I

kDa

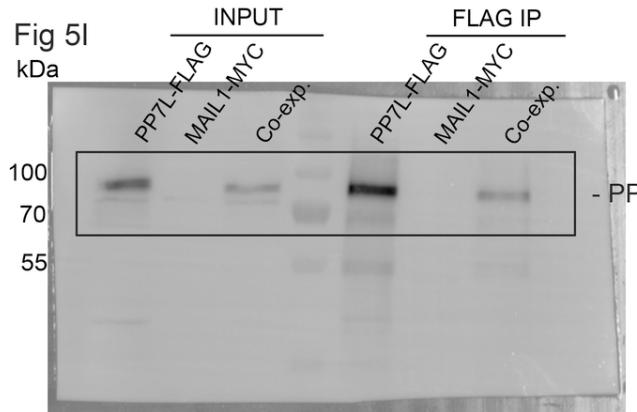


Fig 5I

kDa

